

## **RAW SEQUENCE LISTING**

**The Biotechnology Systems Branch of the Scientific and Technical  
Information Center (STIC) no errors detected.**

Application Serial Number: 10/561,196  
Source: IFWP  
Date Processed by STIC: 09/12/2006

# ***ENTERED***



IFWP

## RAW SEQUENCE LISTING

DATE: 09/12/2006

PATENT APPLICATION: US/10/561,196

TIME: 10:55:08

Input Set : A:\056222-5089-US Sequence Listing.txt

Output Set: N:\CRF4\09122006\J561196.raw

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3 <110> APPLICANT: Dluzewski, Anton R
4   Holder, Anthony A
5   Birdsall, Berry M
6   Babon, Jeff J
7   Martin, Stephen R
8   Morgan, William D
9   Feeney, James
10  Blackman, Michael J
11  Fleck, Suzanne L
12  Saxty, Barbara
14 <120> TITLE OF INVENTION: SCREENING METHOD FOR SUBSTANCES BINDING TO
MEROZOITE SURFACE
15   PROTEIN-1/42
17 <130> FILE REFERENCE: 056222-5089
19 <140> CURRENT APPLICATION NUMBER: US 10/561,196
20 <141> CURRENT FILING DATE: 2005-12-16
22 <150> PRIOR APPLICATION NUMBER: PCT/GB2004/002604
23 <151> PRIOR FILING DATE: 2004-06-18
25 <150> PRIOR APPLICATION NUMBER: GB 0314114.0
26 <151> PRIOR FILING DATE: 2003-06-18
28 <160> NUMBER OF SEQ ID NOS: 15
30 <170> SOFTWARE: PatentIn Ver. 2.1
32 <210> SEQ ID NO: 1
33 <211> LENGTH: 12
34 <212> TYPE: PRT
35 <213> ORGANISM: Plasmodium falciparum
37 <400> SEQUENCE: 1
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43 <210> SEQ ID NO: 2
44 <211> LENGTH: 14
45 <212> TYPE: PRT
46 <213> ORGANISM: Plasmodium vivax
48 <400> SEQUENCE: 2
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54 <210> SEQ ID NO: 3
55 <211> LENGTH: 383
56 <212> TYPE: PRT
57 <213> ORGANISM: Plasmodium falciparum
59 <400> SEQUENCE: 3
60 Gly Glu Ala Val Thr Pro Ser Val Ile Asp Asn Ile Leu Ser Lys Ile
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63 Glu Asn Glu Tyr Glu Val Leu Tyr Leu Lys Pro Leu Ala Gly Val Tyr

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64          20          25          30
66 Arg Ser Leu Lys Lys Gln Leu Glu Asn Asn Val Met Thr Phe Asn Val
67          35          40          45
69 Asn Val Lys Asp Ile Leu Asn Ser Arg Phe Asn Lys Arg Glu Asn Phe
70          50          55          60
72 Lys Asn Val Leu Glu Ser Asp Leu Ile Pro Tyr Lys Asp Leu Thr Ser
73 65          70          75          80
75 Ser Asn Tyr Val Val Lys Asp Pro Tyr Lys Phe Leu Asn Lys Glu Lys
76          85          90          95
78 Arg Asp Lys Phe Leu Ser Ser Tyr Asn Tyr Ile Lys Asp Ser Ile Asp
79          100          105          110
81 Thr Asp Ile Asn Phe Ala Asn Asp Val Leu Gly Tyr Tyr Lys Ile Leu
82          115          120          125
84 Ser Glu Lys Tyr Lys Ser Asp Leu Asp Ser Ile Lys Lys Tyr Ile Asn
85          130          135          140
87 Asp Lys Gln Gly Glu Asn Glu Lys Tyr Leu Pro Phe Leu Asn Asn Ile
88 145          150          155          160
90 Glu Thr Leu Tyr Lys Thr Val Asn Asp Lys Ile Asp Leu Phe Val Ile
91          165          170          175
93 His Leu Glu Ala Lys Val Leu Asn Tyr Thr Tyr Glu Lys Ser Asn Val
94          180          185          190
96 Glu Val Lys Ile Lys Glu Leu Asn Tyr Leu Lys Thr Ile Gln Asp Lys
97          195          200          205
99 Leu Ala Asp Phe Lys Lys Asn Asn Asn Phe Val Gly Ile Ala Asp Leu
100          210          215          220
102 Ser Thr Asp Tyr Asn His Asn Asn Leu Leu Thr Lys Phe Leu Ser Thr
103 225          230          235          240
105 Gly Met Val Phe Glu Asn Leu Ala Lys Thr Val Leu Ser Asn Leu Leu
106          245          250          255
108 Asp Gly Asn Leu Gln Gly Met Leu Asn Ile Ser Gln His Gln Cys Val
109          260          265          270
111 Lys Lys Gln Cys Pro Gln Asn Ser Gly Cys Phe Arg His Leu Asp Glu
112          275          280          285
114 Arg Glu Glu Cys Lys Cys Leu Leu Asn Tyr Lys Gln Glu Gly Asp Lys
115          290          295          300
117 Pro Phe Trp Glu Leu Cys Val Glu Asn Pro Asn Pro Thr Cys Asn Glu
118 305          310          315          320
120 Asn Asn Gly Gly Cys Asp Ala Asp Ala Lys Cys Thr Glu Glu Asp Ser
121          325          330          335
123 Gly Ser Asn Gly Lys Lys Ile Thr Cys Glu Cys Thr Lys Pro Asp Ser
124          340          345          350
126 Tyr Pro Leu Phe Asp Gly Ile Phe Cys Ser Ser Ser Asn Phe Leu Gly
127          355          360          365
129 Ile Ser Phe Leu Leu Ile Leu Met Leu Ile Leu Tyr Ser Phe Ile
130          370          375          380
134 <210> SEQ ID NO: 4
135 <211> LENGTH: 395
136 <212> TYPE: PRT
137 <213> ORGANISM: Plasmodium falciparum

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139 &lt;400&gt; SEQUENCE: 4

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140 Gly Glu Ala Ile Ser Val Thr Met Asp Asn Ile Leu Ser Gly Phe Glu
141   1      5      10      15
143 Asn Glu Tyr Asp Val Ile Tyr Leu Lys Pro Leu Ala Gly Val Tyr Arg
144      20      25      30
146 Ser Leu Lys Lys Gln Ile Glu Lys Asn Ile Ile Thr Phe Asn Leu Asn
147      35      40      45
149 Leu Asn Asp Ile Leu Asn Ser Arg Leu Lys Lys Gly Glu Ala Ile Ser
150      50      55      60
152 Val Thr Met Asp Asn Ile Leu Ser Gly Phe Glu Asn Glu Tyr Asp Val
153   65      70      75      80
155 Ile Tyr Leu Lys Pro Leu Ala Gly Val Tyr Arg Ser Leu Lys Lys Gln
156      85      90      95
158 Ile Glu Lys Asn Ile Ile Thr Phe Asn Leu Asn Leu Asn Asp Ile Leu
159      100     105     110
161 Asn Ser Arg Leu Lys Lys Gly Ile Ser Tyr Tyr Glu Lys Val Leu Ala
162      115     120     125
164 Lys Tyr Lys Asp Asp Leu Glu Ser Ile Lys Lys Val Ile Lys Glu Glu
165      130     135     140
167 Lys Glu Lys Phe Pro Ser Pro Pro Thr Thr Pro Pro Ser Pro Ala
168 145      150      155      160
170 Lys Thr Asp Glu Gln Lys Lys Glu Ser Lys Phe Leu Pro Phe Leu Thr
171      165     170     175
173 Asn Ile Glu Thr Leu Tyr Asn Asn Leu Val Asn Lys Ile Asp Asp Tyr
174      180     185     190
176 Leu Ile Asn Leu Lys Ala Lys Ile Asn Asp Cys Asn Val Glu Lys Asp
177      195     200     205
179 Glu Ala His Val Lys Ile Thr Lys Leu Ser Asp Leu Lys Ala Ile Asp
180      210     215     220
182 Asp Lys Ile Asp Leu Phe Lys Asn Thr Asn Asp Phe Glu Ala Ile Lys
183 225      230      235      240
185 Lys Leu Ile Asn Asp Asp Thr Lys Lys Asp Met Leu Gly Lys Leu Leu
186      245     250     255
188 Ser Thr Gly Leu Val Gln Asn Phe Pro Asn Thr Ile Ile Ser Lys Leu
189      260     265     270
191 Ile Glu Gly Lys Phe Gln Asp Met Leu Asn Ile Ser Gln His Gln Cys
192      275     280     285
194 Val Lys Lys Gln Cys Pro Glu Asn Ser Gly Cys Phe Arg His Leu Asp
195      290     295     300
197 Glu Arg Glu Glu Cys Lys Cys Leu Leu Asn Tyr Lys Gln Glu Gly Asp
198 305      310      315      320
200 Lys Cys Val Glu Asn Pro Asn Pro Thr Cys Asn Glu Asn Asn Gly Gly
201      325     330     335
203 Cys Asp Ala Asp Ala Thr Cys Thr Glu Glu Asp Ser Gly Ser Ser Arg
204      340     345     350
206 Lys Lys Ile Thr Cys Glu Cys Thr Lys Pro Asp Ser Tyr Pro Leu Phe
207      355     360     365
209 Asp Gly Ile Phe Cys Ser Ser Ser Asn Phe Leu Gly Ile Ser Phe Leu
210      370     375     380

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212 Leu Ile Leu Met Leu Ile Leu Tyr Ser Phe Ile
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217 <210> SEQ ID NO: 5
218 <211> LENGTH: 396
219 <212> TYPE: PRT
220 <213> ORGANISM: Plasmodium falciparum
222 <400> SEQUENCE: 5
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224 1 5 10 15
226 Asn Glu Tyr Asp Val Ile Tyr Leu Lys Pro Leu Ala Gly Val Tyr Arg
227 20 25 30
229 Ser Leu Lys Lys Gln Ile Glu Lys Asn Ile Phe Thr Phe Asn Leu Asn
230 35 40 45
232 Leu Asn Asp Ile Leu Asn Ser Arg Leu Lys Lys Arg Lys Tyr Phe Leu
233 50 55 60
235 Asp Val Leu Glu Ser Asp Leu Met Gln Phe Lys His Ile Ser Ser Asn
236 65 70 75 80
238 Glu Tyr Ile Ile Glu Asp Ser Phe Lys Leu Leu Asn Ser Glu Gln Lys
239 85 90 95
241 Asn Thr Leu Leu Lys Ser Tyr Lys Tyr Ile Lys Glu Ser Val Glu Asn
242 100 105 110
244 Asp Ile Lys Phe Ala Gln Glu Gly Ile Ser Tyr Tyr Glu Lys Val Leu
245 115 120 125
247 Ala Lys Tyr Lys Asp Asp Leu Glu Ser Ile Lys Lys Val Ile Lys Glu
248 130 135 140
250 Glu Lys Glu Lys Phe Pro Ser Ser Pro Pro Thr Thr Pro Pro Ser Pro
251 145 150 155 160
253 Ala Lys Thr Asp Glu Gln Lys Lys Glu Ser Lys Phe Leu Pro Phe Leu
254 165 170 175
256 Thr Asn Ile Glu Thr Leu Tyr Asn Asn Leu Val Asn Lys Ile Asp Asp
257 180 185 190
259 Tyr Leu Ile Asn Leu Lys Ala Lys Ile Asn Asp Cys Asn Val Glu Lys
260 195 200 205
262 Asp Glu Ala His Val Lys Ile Thr Lys Leu Ser Asp Leu Lys Ala Ile
263 210 215 220
265 Asp Asp Lys Ile Asp Leu Phe Lys Asn His Asn Asp Phe Glu Ala Ile
266 225 230 235 240
268 Lys Lys Leu Ile Asn Asp Asp Thr Lys Lys Asp Met Leu Gly Lys Leu
269 245 250 255
271 Leu Ser Thr Gly Leu Val Gln Asn Phe Pro Asn Thr Ile Ile Ser Lys
272 260 265 270
274 Leu Ile Glu Gly Lys Phe Gln Asp Met Leu Asn Ile Ser Gln His Gln
275 275 280 285
277 Cys Val Lys Lys Gln Cys Pro Glu Asn Ser Gly Cys Phe Arg His Leu
278 290 295 300
280 Asp Glu Arg Glu Glu Cys Lys Cys Leu Leu Asn Tyr Lys Gln Glu Gly
281 305 310 315 320
283 Asp Lys Cys Val Glu Asn Pro Asn Pro Thr Cys Asn Glu Asn Asn Gly
284 325 330 335

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286 Gly Cys Asp Ala Asp Ala Lys Cys Thr Glu Glu Asp Ser Gly Ser Asn
287           340           345           350
289 Gly Lys Lys Ile Thr Cys Glu Cys Thr Lys Pro Asp Ser Tyr Pro Leu
290           355           360           365
292 Phe Asp Gly Ile Phe Cys Ser Ser Ser Asn Phe Leu Gly Ile Ser Phe
293           370           375           380
295 Leu Leu Ile Leu Met Leu Ile Leu Tyr Ser Phe Ile
296 385           390           395
300 <210> SEQ ID NO: 6
301 <211> LENGTH: 396
302 <212> TYPE: PRT
303 <213> ORGANISM: Plasmodium falciparum
305 <400> SEQUENCE: 6
306 Gly Glu Ala Ile Ser Val Thr Met Asp Asn Ile Leu Ser Gly Phe Glu
307   1           5           10           15
309 Asn Glu Tyr Asp Val Ile Tyr Leu Lys Pro Leu Ala Gly Val Tyr Arg
310           20           25           30
312 Ser Leu Lys Lys Gln Ile Glu Lys Asn Ile Phe Thr Phe Asn Leu Asn
313           35           40           45
315 Leu Asn Asp Ile Leu Asn Ser Arg Leu Lys Lys Arg Lys Tyr Phe Leu
316           50           55           60
318 Asp Val Leu Glu Ser Asp Leu Met Gln Phe Lys His Ile Ser Ser Asn
319 65           70           75           80
321 Glu Tyr Ile Ile Glu Asp Ser Phe Lys Leu Leu Asn Ser Glu Gln Lys
322           85           90           95
324 Asn Thr Leu Leu Lys Ser Tyr Lys Tyr Ile Lys Glu Ser Val Glu Asn
325           100          105          110
327 Asp Ile Lys Phe Ala Gln Glu Gly Ile Ser Tyr Tyr Glu Lys Val Leu
328           115          120          125
330 Ala Lys Tyr Lys Asp Asp Leu Glu Ser Ile Lys Lys Val Ile Lys Glu
331           130          135          140
333 Glu Lys Glu Lys Phe Pro Ser Ser Pro Pro Thr Thr Pro Pro Ser Pro
334 145          150          155          160
336 Ala Lys Thr Asp Glu Gln Lys Lys Glu Ser Lys Phe Leu Pro Phe Leu
337           165          170          175
339 Thr Asn Ile Glu Thr Leu Tyr Asn Asn Leu Val Asn Lys Ile Asp Asp
340           180          185          190
342 Tyr Leu Ile Asn Leu Lys Ala Lys Ile Asn Asp Cys Asn Val Glu Lys
343           195          200          205
345 Asp Glu Ala His Val Lys Ile Thr Lys Leu Ser Asp Leu Lys Ala Ile
346           210          215          220
348 Asp Asp Lys Ile Asp Leu Phe Lys Asn His Asn Asp Phe Asp Ala Ile
349 225          230          235          240
351 Lys Lys Leu Ile Asn Asp Asp Thr Lys Lys Asp Met Leu Gly Lys Leu
352           245          250          255
354 Leu Ser Thr Gly Leu Val Gln Asn Phe Pro Asn Thr Ile Ile Ser Lys
355           260          265          270
357 Leu Ile Glu Gly Lys Phe Gln Asp Met Leu Asn Ile Ser Gln His Gln
358           275          280          285

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VERIFICATION SUMMARY

PATENT APPLICATION: US/10/561,196

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